

Scientific Programming

Lecture A07 – Pandas

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Documentation

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What is Pandas?

Pandas

A freely available library for loading, manipulating, and visualizing sequential and tabular data, such as time series or micro-arrays.

Features

- Loading and saving with “standard” tabular file formats:
 - CSV (Comma-separated Values)
 - TSV (Tab-separated Values)
 - Excel files
 - Database formats, etc.
- Flexible indexing and aggregation of series and tables
- Efficient numerical/statistical operations (e.g. broadcasting)
- Pretty, straightforward visualization

Some links

Official Pandas website

<http://pandas.pydata.org/>

Official documentation

<http://pandas.pydata.org/pandas-docs/stable/dsintro.html>

Source code

<https://github.com/pandas-dev/pandas/>

A short demonstration – Iris Dataset

```
SepalLength,SepalWidth,PetalLength,PetalWidth,Name
```

```
5.1,3.5,1.4,0.2,Iris-setosa
```

```
4.9,3.0,1.4,0.2,Iris-setosa
```

```
...
```

```
5.0,3.3,1.4,0.2,Iris-setosa
```

```
7.0,3.2,4.7,1.4,Iris-versicolor
```

```
6.4,3.2,4.5,1.5,Iris-versicolor
```

```
...
```

```
5.7,2.8,4.1,1.3,Iris-versicolor
```

```
6.3,3.3,6.0,2.5,Iris-virginica
```

```
5.8,2.7,5.1,1.9,Iris-virginica
```

```
...
```

```
https://drive.google.com/open?id=0B0wILN942aEVYTVBekRHLTNON3c
```

```
https://en.wikipedia.org/wiki/Iris\_flower\_data\_set
```

A short demonstration – Iris Dataset

In an effort to understand the dataset, we would like to visualize the relation between the four properties for the case of Iris virginica.

- Load the dataset by parsing all the rows in the file
- Keep only the rows pertaining to Iris virginica
- Compute statistics on the values of the rows, making sure to convert from strings to float's as required
- Actually draw the plots by using a specialized plotting library.

A short demonstration – Iris Dataset

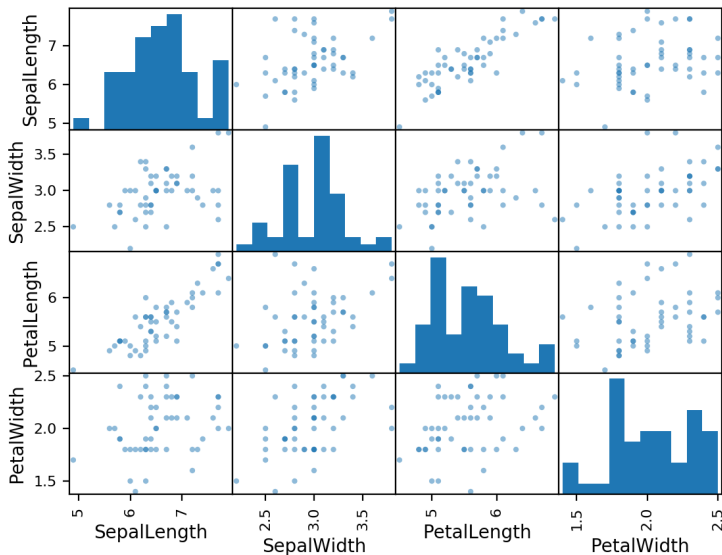
```
import pandas as pd
from pandas.plotting import scatter_matrix
import matplotlib.pyplot as plt

df = pd.read_csv("iris.csv")

scatter_matrix(df[df.Name == "Iris-virginica"])

plt.show()
```

A short demonstration – Iris Dataset



Introduction to Pandas

Pandas provides a couple of very useful datatypes:

- **Series** represents 1D data, like time series, calendars, the output of one-variable functions, etc.
- **DataFrame** represents 2D data, like a column-separated-values (CSV) file, a microarray, a database table, a matrix, etc.

Each column of a **DataFrame** is a **Series**.

- That's why we will see how the **Series** data type works first.
- Most of what we will say about **Series** also applies to **DataFrames**.

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Pandas: Series

Series

A **Series** is a **one-dimensional array** with a **labeled axis**, that can hold arbitrary objects.

The axis is called the **index**, and can be used to access the elements; it is very flexible, and not necessarily numerical.

It works partially like a **list** and partially like a **dict**.

Creating a Series (1)

It is possible to specify just the series data, associating an **implicit** numeric index.

```
import pandas as pd
s = pd.Series(["a", "b", "c"])
print(s)
```

```
0    a
1    b
2    c
dtype: object
```

Creating a Series (2)

It is possible to specify both the series data and the **explicit** index, separately:

```
import pandas as pd
s = pd.Series(["a", "b", "c"], index=[2, 5, 8])
print(s)
```

```
2    a
```

```
5    b
```

```
8    c
```

```
dtype: object
```

Creating a Series (3)

It is possible to specify both the series data and the index, as a single dictionary:

```
import pandas as pd
s = pd.Series({"a": "A", "b": "B", "c": "C"})
print(s)
```

```
a    A
```

```
b    B
```

```
c    C
```

```
dtype: object
```

Creating a Series (4)

If given a single scalar (e.g. an integer), the series constructor will replicate it for all indices (that need to be specified)

```
import pandas as pd
s = pd.Series(3, index=range(5))
print(s)
```

```
0    3
```

```
1    3
```

```
2    3
```

```
3    3
```

```
4    3
```

```
dtype: int64
```

Accessing a Series

Let's create a Series representing the hours of sleep we had the chance to get each day of the past week. We may now access it through either the position (as a list) or the index (as a dict)

```
import pandas as pd
days = ["mon", "tue", "wed", "thu", "fri"]
sleephours = [6, 2, 8, 5, 9]
s = pd.Series(sleephours, index=days)
print(s["mon"])
s["tue"]=3
print(s[1])
```

6

3

Accessing a Series

- If a label is not contained, an exception is raised.
- Using the get method, a missing label will return None or specified default

```
import pandas as pd
days = ["mon", "tue", "wed", "thu", "fri"]
sleephours = [6, 2, 8, 5, 9]
s = pd.Series(sleephours, index=days)
```

```
print(s["sat"])
print(s.get('sat'))
```

```
KeyError: 'sat'
None
```

Slicing a Series

We can also slice the positions, like we would do with a list. Note that both the data and the index are extracted correctly. It also works with labels.

```
print(s[-3:])  
print(s["tue":"thu"])
```

```
wed    8  
thu    5  
fri    9  
dtype: int64  
tue    2  
wed    8  
thu    5  
dtype: int64
```

Head and tail

The first and last `n` elements can be extracted also using `head()` and `tail()`.

```
print(s.head(2))  
print(s.tail(3))
```

```
mon    6  
tue    2  
dtype: int64
```

```
wed    8  
thu    5  
fri    9  
dtype: int64
```

List of indexes

You can also explicitly pass a list of positions. Tuples do not work, because they are interpreted as potential indexes.

```
print(s[[0, 1, 2]])  
print(s[["mon", "wed", "fri"]])
```

```
mon    6  
tue    2  
wed    8  
dtype: int64
```

```
mon    6  
wed    8  
fri    9  
dtype: int64
```

Operator broadcasting

The **Series** class automatically broadcasts arithmetical operations by a scalar to all of the elements.

```
print(s)
```

```
mon    6  
tue    2  
wed    8  
thu    5  
fri    9  
dtype: int64
```

```
print(s+1)
```

```
mon    7  
tue    3  
wed    9  
thu    6  
fri   10  
dtype: int64
```

```
print(s*2)
```

```
mon   12  
tue    4  
wed   16  
thu   10  
fri   18  
dtype: int64
```

Note

The concept of operator broadcasting was taken from the `numpy` library, and is one of the key features for writing efficient, clean numerical code in Python.

In a way, it is a “generalized” version of scalar products (from linear algebra).

The rules governing how broadcasting is applied can be pretty complex (and confusing). For the moment, we will cover constant broadcasting only.

Masking and filtering

- Besides numerical operators, we can apply boolean conditions. The result is called a **mask**.
- Masks can be used to **filter** the elements of a **Series** according to a given condition.

```
print(s)
```

```
mon    6
tue    2
wed    8
thu    5
fri    9
dtype: int64
```

```
print(s>=6)
```

```
mon    True
tue    False
wed    True
thu    False
fri    True
dtype: bool
```

```
print(s[s>=6])
```

```
mon    6
wed    8
fri    9
dtype: int64
```

Automatic label assignments

Operations between multiple time series are automatically aligned by label, meaning that elements with the same label are matched prior to carrying out the operation.

```
print(s[1:])
```

```
tue    2
wed    8
thu    5
fri    9
dtype: int64
```

```
print(s[:-1])
```

```
mon    6
tue    2
wed    8
thu    5
dtype: int64
```

```
print(s[1:]+s[:-1])
```

```
fri    NaN
mon    NaN
thu    10.0
tue    4.0
wed    16.0
dtype: float64
```


Not-a-Number (NaN)

The index of the resulting **Series** is the union of the indices of the operands. What happens depend on whether a given label appears in both input Series or not:

- For common labels (in our case "tue", "wed", "thu"), the output **Series** contains the sum of the aligned elements.
- For labels appearing in only one of the operands ("mon" and "fri"), the result is a NaN, i.e. not-a-number.

NaN is just a symbolic constant that specifies that the object is a number-like entity with an invalid or undefined value.

Dealing with missing values

There are different strategies for dealing with nan's. There is no "best" strategy: you have to pick one depending on the problem you are trying to solve.

```
t = s[1:]+s[:-1]
print(t)
```

```
fri    NaN
mon    NaN
thu    10.0
tue     4.0
wed    16.0
dtype: float64
```

```
nt = t.dropna()
print(nt)
```

```
thu    12.0
tue     6.0
wed    18.0
dtype: float64
```

```
zt = t.fillna(0.0)
print(zt)
```

```
fri    0.0
mon    0.0
thu    12.0
tue     6.0
wed    18.0
dtype: float64
```

Automatic label assignments

Through the method `add`, it is possible to assign a fill value to the missing entries of the series to be added, in order to get a real sum.

```
print(s[1:])
```

```
tue    2
wed    8
thu    5
fri    9
dtype: int64
```

```
print(s[:-1])
```

```
mon    6
tue    2
wed    8
thu    5
dtype: int64
```

```
print(s[1:].add(s[:-1],
               fill_value=0))
```

```
fri    9.0
mon    6.0
thu   10.0
tue    4.0
wed   16.0
dtype: float64
```

Computing statistics

```

print(s)
mon      6
tue      2
wed      8
thu      5
fri      9
dtype: int64

print(s.sum())      30
print(s.prod())    4320
print(s.max())      9
print(s.argmax())   fri
print(s.mean())     6.0
print(s.var())      7.5
print(s.std())      2.7386127875258306
print(s.median())   6.0

print(s.cumsum())
mon      6
tue      8
wed     16
thu     21
fri     30
dtype: int64

```

Computing statistics

```

print(s)
mon    6
tue    2
wed    8
thu    5
fri    9
dtype: int64

print(s.quantile(0.5))
6.0

print(s.quantile(
    [0.25, 0.5, 0.75]
))
0.25    5.0
0.50    6.0
0.75    8.0
dtype: float64

print(s[s >=
    s.quantile(0.5)
])
mon    6
wed    8
fri    9
dtype: int64

```

Computing statistics

```
print(s)                # Pearson corr.
                        print(s.corr(s))                1.0
mon      6
tue      2                # Spearman corr.
wed      8                print(s.corr(s,                1.0
thu      5                method="spearman"
fri      9                )
dtype: int64

                        # Autocorrelation
                        # with time lag
                        print(s.autocorr(lag=0))        1.0
                        print(s.autocorr(lag=1))        -0.548128127763
                        print(s.autocorr(lag=2))        0.995870594886
```

Series: Conclusion

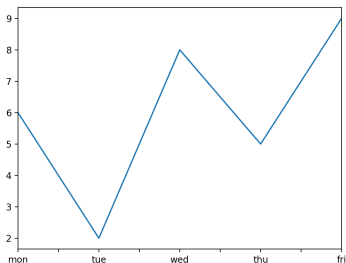
A quick way to get several useful statistics is to use `s.describe()`. Anyway, the list of statistical methods associated with `Series` is larger than this.

```
print(s.describe())
```

```
count    5.000000
mean     6.000000
std      2.738613
min      2.000000
25%      5.000000
50%      6.000000
75%      8.000000
max      9.000000
dtype: float64
```

Series: Plotting

```
import pandas as pd
import matplotlib.pyplot as plt
sleephours = [6, 2, 8, 5, 9]
days = ["mon", "tue", "wed", "thu", "fri"]
s = pd.Series(sleephours, index=days)
s.plot()
plt.show()
```



```
import pandas as pd
import matplotlib.pyplot as plt
sleephours = [6, 2, 8, 5, 9]
days = ["mon", "tue", "wed", "thu", "fri"]
s = pd.Series(sleephours, index=days)
s.plot(kind="bar")
plt.show()
```

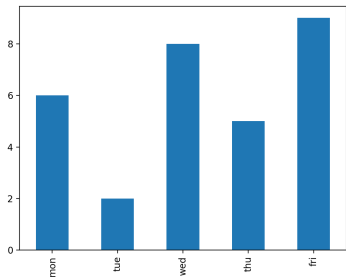


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DataFrame

Pandas **DataFrame** is the 2D analogue of a **Series**: it is essentially a table of heterogeneous objects.

A **DataFrame** holds three major attributes:

- the **index**, which holds the labels of the rows
- the **columns**, which holds the labels of the columns
- the **shape**, which describes the dimension of the table

When you extract a column from a **DataFrame** you get a proper **Series**, and you can operate on it using all the tools presented in the previous section.

Further, most (not all) of the operations that you can do on a **Series**, you can also do on an entire **DataFrame**.

Creating a DataFrame from a dictionary of Series

```
import pandas as pd
d = { "name": pd.Series(["bobby", "ronald", "ronald", "ronald"]),
      "surname": pd.Series(["fisher", "fisher", "reagan", "mcdonald"]) }
df = pd.DataFrame(d)
print(df)
print(df.columns)
print(df.index)
print(df.shape)
```

- The keys of the dictionary became the columns of the dataframe
- The index of the various Series became the index of the dataframe.

```
      name  surname
0  bobby   fisher
1  ronald   fisher
2  ronald   reagan
3  ronald  mcdonald
Index(['name', 'surname'], dtype='object')
RangeIndex(start=0, stop=4, step=1)
(4, 2)
```

Creating a DataFrame from a dictionary of Series

```
import pandas as pd
d = { "x": pd.Series([0, 0], index=["a", "b"]),
      "y": pd.Series([0, 0], index=["b", "c"])}
df = pd.DataFrame(d)
print(df)
print(df.columns)
print(df.index)
print(df.shape)
```

- If the index of the input Series do not match, since label alignment applies, the missing values are treated as NaN's.

```
      x    y
a  0.0  NaN
b  0.0  0.0
c  NaN  0.0
Index(['x', 'y'], dtype='object')
Index(['a', 'b', 'c'], dtype='object')
(3,2)
```

Creating a DataFrame from a dictionary of lists

```
import pandas as pd
d = { "column1": [1., 2., 6., -1.],
      "column2": [0., 1., -2., 4.] }
df = pd.DataFrame(d)
print(df)
print(df.columns)
print(df.index)
print(df.shape)
```

- The columns are taken from the keys
- The index is set to the default one

```
   column1  column2
0      1.0      0.0
1      2.0      1.0
2      6.0     -2.0
3     -1.0      4.0
Index(['column1', 'column2'], dtype='object')
RangeIndex(start=0, stop=4, step=1)
(4,2)
```

Creating a DataFrame from a dictionary of lists, with index

```
import pandas as pd
d = { "column1": [1., 2., 6., -1.],
      "column2": [0., 1., -2., 4.] }
df = pd.DataFrame(d, index=["a", "b", "c", "d"])
print(df)
print(df.columns)
print(df.index)
print(df.shape)
```

- A custom index can be specified the usual way

```

   column1  column2
a         1.0       0.0
b         2.0       1.0
c         6.0      -2.0
d        -1.0       4.0
Index(['column1', 'column2'], dtype='object')
RangeIndex(start=0, stop=4, step=1)
(4,2)
```

Creating a DataFrame from a list of dictionaries

```
import pandas as pd
d = [ {"a": 1, "b": 2},
      {"a": 2, "c": 3},
]
df = pd.DataFrame(d)
print(df)
print(df.columns)
print(df.index)
print(df.shape)
```

```
   a    b    c
0  1  2.0 NaN
1  2  NaN  3.0
Index(['a', 'b', 'c'], dtype='object')
RangeIndex(start=0, stop=2, step=1)
(2,3)
```

- The columns are taken from the keys of the dictionaries
- The index is the default one.
- Since not all common keys appear in all input dictionaries, missing values (i.e. NaN's) are automatically added.

Loading a CSV file

```
import pandas as pd
df = pd.read_csv("iris.csv")
print(df.columns)
print(df.index)
print(df.shape)
```

```
Index(['SepalLength', 'SepalWidth', 'PetalLength', 'PetalWidth', 'Name'],
      dtype='object')
```

```
RangeIndex(start=0, stop=150, step=1)
(150, 5)
```

| | SepalLength | SepalWidth | PetalLength | PetalWidth | Name |
|---|-------------|------------|-------------|------------|-------------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 | Iris-setosa |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 | Iris-setosa |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 | Iris-setosa |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 | Iris-setosa |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 | Iris-setosa |
| 5 | 5.4 | 3.9 | 1.7 | 0.4 | Iris-setosa |

...

help(pd.read_csv)

Help on function read_csv in module pandas.io.parsers:

```
read_csv(filepath_or_buffer, sep=',', delimiter=None, header='infer',
names=None, index_col=None, usecols=None, squeeze=False, prefix=None,
mangle_dupe_cols=True, dtype=None, engine=None, converters=None,
true_values=None, false_values=None, skipinitialspace=False,
skiprows=None, nrows=None, na_values=None, keep_default_na=True,
na_filter=True, verbose=False, skip_blank_lines=True, parse_dates=False,
infer_datetime_format=False, keep_date_col=False, date_parser=None,
dayfirst=False, iterator=False, chunksize=None, compression='infer',
thousands=None, decimal=b'.' , lineterminator=None, quotechar='"',
quoting=0, escapechar=None, comment=None, encoding=None, dialect=None,
tupleize_cols=False, error_bad_lines=True, warn_bad_lines=True,
skipfooter=0, skip_footer=0, doublequote=True, delim_whitespace=False,
as_recarray=False, compact_ints=False, use_unsigned=False,
low_memory=True, buffer_lines=None, memory_map=False, float_precision=None)
```

Read CSV (comma-separated) file into DataFrame

Loading a malformed Tab-Separated File

<https://drive.google.com/open?id=0B0wILN942aEVeWdScy1wQnA3LTA>

It describes a mapping from UniProt protein IDs (i.e. sequences) to hits in the Protein Data Bank (i.e. structures). The TSV file looks like this:

```
# 2014/07/08 - 14:59
```

```
SP_PRIMARY    PDB
```

```
A0A011        3vk5;3vka;3vkb;3vkc;3vkd
```

```
A0A2Y1        2jrd
```

```
A0A585        4mnq
```

```
A0A5B4        2ij0
```

```
A0A5B9        2bnq;2eyr;2eys;2eyt;3kxf;3o6f;3o8x;3o9w;3qux;3t0e;3
```

```
A0A5E3        1hq4;1ob1
```

```
A0AEF5        4iu2;4iu3
```

```
A0AEF6        4iu2;4iu3
```

```
A0A007        2ib5
```

Loading a malformed Tab-Separated File

We can use the `sep` (separator) argument of the `read_csv()` method to take care of the TABs.

```
df = pd.read_csv("uniprot_pdb.tsv", sep="\t")
print(df.shape)
print(df.columns)
print(df.head(3))
```

```
(33636, 1)
```

```
Index(['# 2014/07/08 - 14:59'], dtype='object')
```

```
      # 2014/07/08 - 14:59
```

```
SP_PRIMARY          PDB
```

```
A0A011      3vk5;3vka;3vkb;3vkc;3vkd
```

```
A0A2Y1          2jrd
```

Problem: the first line contains only one column, so Pandas think that there is only one column

Loading a malformed Tab-Separated File

Argument `skiprows` is used to skip the first line, which is a comment.

```
df = pd.read_csv("uniprot_pdb.tsv", sep="\t", skiprows=1)
print(df.shape)
print(df.columns)
print(df.head(3))
```

```
(33636, 2)
```

```
Index(['SP_PRIMARY', 'PDB'], dtype='object')
```

| | SP_PRIMARY | PDB |
|---|------------|--------------------------|
| 0 | AOA011 | 3vk5;3vka;3vkb;3vkc;3vkd |
| 1 | AOA2Y1 | 2jrd |
| 2 | AOA585 | 4mnq |

Loading a malformed Tab-Separated File

Argument `comment` is used to skip all the lines that start with #

```
df = pd.read_csv("uniprot_pdb.tsv", sep="\t", comment="#")
print(df.shape)
print(df.columns)
print(df.head(3))
```

```
(33636, 2)
```

```
Index(['SP_PRIMARY', 'PDB'], dtype='object')
```

| | SP_PRIMARY | PDB |
|---|------------|--------------------------|
| 0 | AOA011 | 3vk5;3vka;3vkb;3vkc;3vkd |
| 1 | AOA2Y1 | 2jrd |
| 2 | AOA585 | 4mnq |

Extracting rows and columns

| Operation | Syntax | Result |
|--------------------------------|------------------------------|------------------|
| Select column | <code>df[col]</code> | Series |
| Select multiple columns | <code>df[[col1,col2]]</code> | DataFrame |
| Select row by label | <code>df.loc[label]</code> | Series |
| Select row by integer location | <code>df.iloc[loc]</code> | Series |
| Slice rows | <code>df[5:10]</code> | DataFrame |
| Select rows by boolean vector | <code>df[bool_vec]</code> | DataFrame |

Extracting a subset of the Iris Dataset

For simplicity, in the following examples we will use a random sample taken from the iris dataset, computed like this

```
import numpy as np
import pandas as pd
np.random.seed(0)
df = pd.read_csv("iris.csv")
small = df.iloc[np.random.permutation(df.shape[0])].head()
print(small.shape)
print(small)
```

Brief explanation: here we use `numpy.random.permutation()` to generate a random permutation of the indices from 0 to `df.shape[0]`, i.e. the number of rows in the Iris dataset; then we use this permutation as row indices to permute all the rows in `df`; finally, we take the first 5 rows of the permuted `df` using the `head()` method.

Extracting a subset of the Iris Dataset

(5, 5)

| | SepalLength | SepalWidth | PetalLength | PetalWidth | Name |
|-----|-------------|------------|-------------|------------|------------------|
| 114 | 5.8 | 2.8 | 5.1 | 2.4 | Iris-virginica |
| 62 | 6.0 | 2.2 | 4.0 | 1.0 | Iris-versicolour |
| 33 | 5.5 | 4.2 | 1.4 | 0.2 | Iris-setosa |
| 107 | 7.3 | 2.9 | 6.3 | 1.8 | Iris-virginica |
| 7 | 5.0 | 3.4 | 1.5 | 0.2 | Iris-setosa |

Extracting a column

It is possible to access the columns of `small` with the `[]` notation. The result is a `Series`.

If the name of the column is compatible with the Python conventions for variable names, you can also treat columns as if they were actual attributes of the dataframes.

```
print(small["Name"])      OR      print(small.Name)
```

```
114    Iris-virginica
62     Iris-versicolor
33      Iris-setosa
107    Iris-virginica
7      Iris-setosa
Name: Name, dtype: object
```

Extracting multiple columns

It is possible to extract multiple columns in one go, by specifying a list of columns; the result is a `DataFrame`

```
print(small[["SepalLength", "PetalLength"]])
```

| | SepalLength | PetalLength |
|-----|-------------|-------------|
| 114 | 5.8 | 5.1 |
| 62 | 6.0 | 4.0 |
| 33 | 5.5 | 1.4 |
| 107 | 7.3 | 6.3 |
| 7 | 5.0 | 1.5 |

Extracting a row

To extract a row, it is possible to use the `loc` and `iloc` attributes by specifying a label or a position, respectively.

The result is a **Series**

```
print(small.loc[114])           OR           print(small.iloc[0])
```

```
SepalLength      5.8
SepalWidth        2.8
PetalLength       5.1
PetalWidth        2.4
Name              Iris-virginica
Name: 114, dtype: object
```

Extracting multiple rows

To extract multiple rows, it is possible to use the `loc` and `iloc` attributes by specifying a list of labels or positions.

The result is a `Dataframe`

```
print(small.loc[[114,62,33]]) OR print(small.iloc[[0,1,2]])
```

| | SepalLength | SepalWidth | PetalLength | PetalWidth | Name |
|-----|-------------|------------|-------------|------------|-----------------|
| 114 | 5.8 | 2.8 | 5.1 | 2.4 | Iris-virginica |
| 62 | 6.0 | 2.2 | 4.0 | 1.0 | Iris-versicolor |
| 33 | 5.5 | 4.2 | 1.4 | 0.2 | Iris-setosa |

Broadcasting

Broadcasting is applied automatically to all rows, or to the entire table.

```
print(small["SepalLength"] + small["SepalWidth"])
print(small+small)
```

```
114      8.6
62       8.2
33       9.7
107     10.2
7        8.4
dtype: float64
```

| | SepalLength | SepalWidth | PetalLength | PetalWidth | |
|-----|-------------|------------|-------------|------------|-----------------------|
| 114 | 11.6 | 5.6 | 10.2 | 4.8 | Iris-virginicaIris- |
| 62 | 12.0 | 4.4 | 8.0 | 2.0 | Iris-versicolorIris-v |
| 33 | 11.0 | 8.4 | 2.8 | 0.4 | Iris-setosaIris- |
| 107 | 14.6 | 5.8 | 12.6 | 3.6 | Iris-virginicaIris- |
| 7 | 10.0 | 6.8 | 3.0 | 0.4 | Iris-setosaIris- |

Masking

Masking works as well.

```
print(small["PetalLength"][small.PetalLength > 5])
print(small["Name"][small.Name == "Iris-virginica"])
print(small[["Name", "PetalLength", "SepalLength"][
    small.Name == "Iris-virginica"]])
```

```
114    5.1
107    6.3
Name: PetalLength, dtype: float64
```

```
114    Iris-virginica
107    Iris-virginica
Name: Name, dtype: object
```

| | Name | PetalLength | SepalLength |
|-----|----------------|-------------|-------------|
| 114 | Iris-virginica | 5.1 | 5.8 |
| 107 | Iris-virginica | 6.3 | 7.3 |

Statistics

Statistics can be computed on rows, columns, or the whole table.

```
print(small.loc[114][:-1].mean()) # Exclude last column, Name
print(small.PetalLength.mean())
print(small.mean())
```

```
4.025
```

```
3.66
```

```
SepalLength    5.92
SepalWidth     3.10
PetalLength    3.66
PetalWidth     1.12
dtype: float64
```

All together!

```
print(small[["Name", "PetalLength", "SepalLength"]][
    small.PetalLength > small.PetalLength.mean()])
```

| | Name | PetalLength | SepalLength |
|-----|-----------------|-------------|-------------|
| 114 | Iris-virginica | 5.1 | 5.8 |
| 62 | Iris-versicolor | 4.0 | 6.0 |
| 107 | Iris-virginica | 6.3 | 7.3 |

Merge

Merging different dataframes is performed using the `merge()` function.

Merging means that, given two tables with a common column name, first the rows with the same column value are matched; then a new table is created by concatenating the matching rows.

```
sequences = pd.DataFrame({
    "id": ["Q99697", "O18400", "P78337", "Q9W5Z2"],
    "seq": ["METNCR", "MDRSSA", "MDAFKG", "MTSMKD"],
})

names = pd.DataFrame({
    "id": ["Q99697", "O18400", "P78337", "P59583"],
    "name": ["PITX2_HUMAN", "PITX_DROME",
            "PITX1_HUMAN", "WRK32_ARATH"],
})
```

Merge - Inner

```
print(sequences)
print(names)
print(pd.merge(sequences, names, on="id", how="inner"))
```

| | id | seq | | id | name |
|---|--------|---------|---|--------|-------------|
| 0 | Q99697 | METNCR | 0 | Q99697 | PITX2_HUMAN |
| 1 | O18400 | MDR SSA | 1 | O18400 | PITX_DROME |
| 2 | P78337 | MDAFKG | 2 | P78337 | PITX1_HUMAN |
| 3 | Q9W5Z2 | MTSMKD | 3 | P59583 | WRK32_ARATH |

| | id | seq | name |
|---|--------|---------|-------------|
| 0 | Q99697 | METNCR | PITX2_HUMAN |
| 1 | O18400 | MDR SSA | PITX_DROME |
| 2 | P78337 | MDAFKG | PITX1_HUMAN |

Mismatched ids are
dropped

Merge - Left

```
print(sequences)
print(names)
print(pd.merge(sequences, names, on="id", how="left"))
```

| | id | seq | | id | name |
|---|--------|--------|---|--------|-------------|
| 0 | Q99697 | METNCR | 0 | Q99697 | PITX2_HUMAN |
| 1 | O18400 | MDR5SA | 1 | O18400 | PITX_DROME |
| 2 | P78337 | MDAFKG | 2 | P78337 | PITX1_HUMAN |
| 3 | Q9W5Z2 | MTSMKD | 3 | P59583 | WRK32_ARATH |

| | id | seq | name |
|---|--------|--------|-------------|
| 0 | Q99697 | METNCR | PITX2_HUMAN |
| 1 | O18400 | MDR5SA | PITX_DROME |
| 2 | P78337 | MDAFKG | PITX1_HUMAN |
| 3 | Q9W5Z2 | MTSMKD | NaN |

The ids are taken
from the left table

Merge - Right

```
print(sequences)
print(names)
print(pd.merge(sequences, names, on="id", how="right"))
```

| | id | seq | | id | name |
|---|--------|---------|---|--------|-------------|
| 0 | Q99697 | METNCR | 0 | Q99697 | PITX2_HUMAN |
| 1 | O18400 | MDR SSA | 1 | O18400 | PITX_DROME |
| 2 | P78337 | MDAFKG | 2 | P78337 | PITX1_HUMAN |
| 3 | Q9W5Z2 | MTSMKD | 3 | P59583 | WRK32_ARATH |

| | id | seq | name |
|---|--------|---------|-------------|
| 0 | Q99697 | METNCR | PITX2_HUMAN |
| 1 | O18400 | MDR SSA | PITX_DROME |
| 2 | P78337 | MDAFKG | PITX1_HUMAN |
| 3 | P59583 | NaN | WRK32_ARATH |

The ids are taken
from the right table

Merge - Outer

```
print(sequences)
print(names)
print(pd.merge(sequences, names, on="id", how="outer"))
```

| | id | seq | | id | name |
|---|--------|--------|---|--------|-------------|
| 0 | Q99697 | METNCR | 0 | Q99697 | PITX2_HUMAN |
| 1 | O18400 | MDR5SA | 1 | O18400 | PITX_DROME |
| 2 | P78337 | MDAFKG | 2 | P78337 | PITX1_HUMAN |
| 3 | Q9W5Z2 | MTSMKD | 3 | P59583 | WRK32_ARATH |

| | id | seq | name |
|---|--------|--------|-------------|
| 0 | Q99697 | METNCR | PITX2_HUMAN |
| 1 | O18400 | MDR5SA | PITX_DROME |
| 2 | P78337 | MDAFKG | PITX1_HUMAN |
| 3 | Q9W5Z2 | MTSMKD | NaN |
| 4 | P59583 | NaN | WRK32_ARATH |

All ids are retained

Grouping Tables

The `groupby` method is essential for efficiently performing operations on groups of rows.

Given the Iris dataset, we want to compute the average of the four columns for each of the three different Iris species.

The result should be a dataframe with 3 species (rows) by 4 columns (petal/sepal length/width).

Grouping Tables

```
import pandas as pd
iris = pd.read_csv("iris.csv")
print(iris.info())
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
SepalLength    150 non-null float64
SepalWidth     150 non-null float64
PetalLength    150 non-null float64
PetalWidth     150 non-null float64
Name           150 non-null object
dtypes: float64(4), object(1)
memory usage: 5.9+ KB
```

Grouping Tables

Iterating over the grouped variable returns (value-of-Name, DataFrame) tuples:

- The 1st item is the value of the column **Name** shared by the group
- The 2nd item is a **DataFrame** including only the rows in that group.

```
grouped = iris.groupby(iris.Name)
for group in grouped:
    print(group[0], group[1].shape)
```

```
Iris-setosa (50, 5)
```

```
Iris-versicolor (50, 5)
```

```
Iris-virginica (50, 5)
```


Grouping Tables

It is possible to apply some transformation (e.g. `mean()`) to the individual groups automatically, using the `aggregate()` method directly on the grouped variable. The result of `aggregate()` is a dataframe.

```
iris_mean_by_name = grouped.aggregate(pd.DataFrame.mean)
print(iris_mean_by_name)
```

| Name | SepalLength | SepalWidth | PetalLength | PetalWidth |
|-----------------|-------------|------------|-------------|------------|
| Iris-setosa | 5.006 | 3.418 | 1.464 | 0.244 |
| Iris-versicolor | 5.936 | 2.770 | 4.260 | 1.326 |
| Iris-virginica | 6.588 | 2.974 | 5.552 | 2.026 |